

Acceleration Data Structures for Ray Tracing

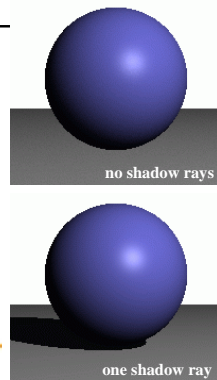
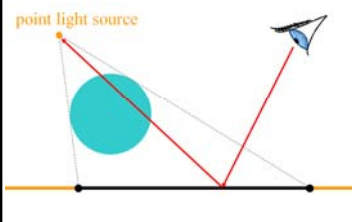
Thanks to Fredo Durand and Barb Cutler

Ray-tracing Acceleration

- Motivation – Distribution Ray Tracing
 - Soft shadows
 - Antialiasing (getting rid of jaggies)
 - Glossy reflection
 - Motion blur
 - Depth of field (focus)
- Bounding Boxes
- Spatial Acceleration Data Structures
- Flattening the Transformation Hierarchy

Shadows

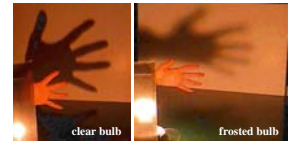
- one shadow ray per intersection per point light source



Shadows & Light Sources



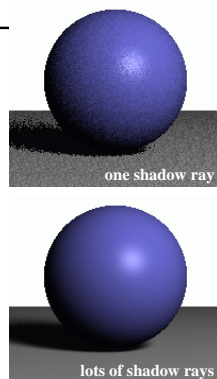
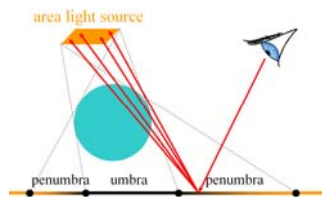
<http://www.davidfay.com/index.php>



<http://www.pa.uky.edu/~scworks/light/preview/bulb2.htm>

Soft Shadows

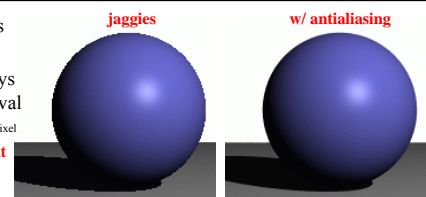
- Multiple shadow rays to sample area light source
- Distribute rays over an interval – light source



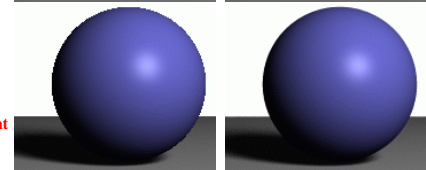
Antialiasing – Supersampling

- Multiple rays per pixel
- Distribute rays over an interval – multiple rays per pixel

point light



area light



Reflection

- one reflection ray per intersection

perfect mirror

Glossy Reflection

- multiple reflection rays
- Distribute rays over an interval – multiple reflection rays

polished surface

Justin Legakis

Motion Blur

- Sample objects temporally
- Distribute rays over an interval - time

Rob Cook

Depth of Field

- Distribute rays over an interval - lens area

Justin Legakis

Ray Tracing Algorithm Analysis

- Ray casting
- Lots of primitives
- Recursive
- Distributed Ray Tracing Effects
 - Soft shadows
 - Anti-aliasing
 - Glossy reflection
 - Motion blur
 - Depth of field

cost \approx height * width * num primitives * intersection cost * size of recursive ray tree * num shadow rays * num supersamples * num glossy rays * num temporal samples * num focal samples * ...

can we reduce this?

Ray-Tracing Taxonomy

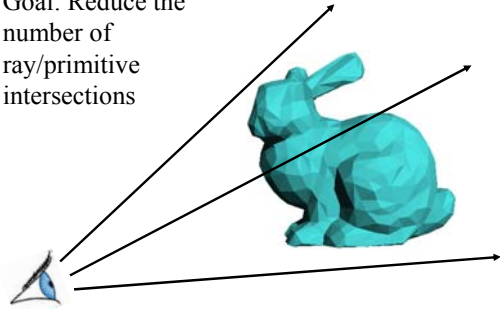
Figure 1: A broad classification of acceleration techniques.
by James Arvo and David Kirk

Ray-tracing Acceleration

- Motivation – Distribution Ray Tracing
- **Bounding Boxes**
 - of each primitive
 - of groups
 - of transformed primitives
- Spatial Acceleration Data Structures
- Flattening the Transformation Hierarchy

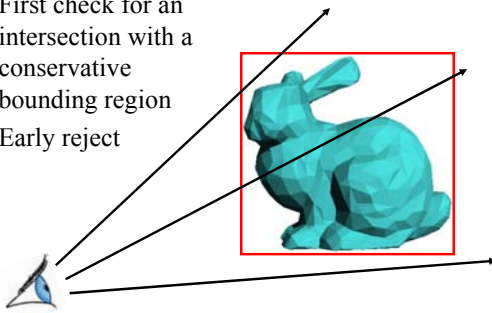
Acceleration of Ray Casting

- Goal: Reduce the number of ray/primitive intersections



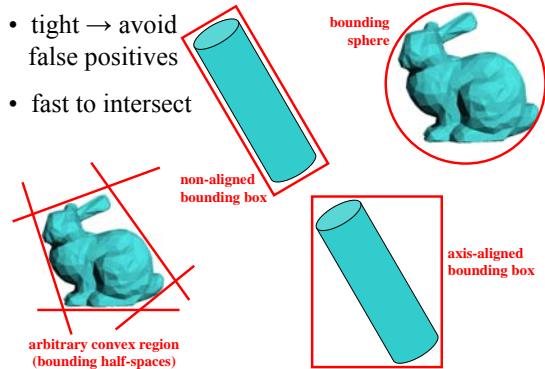
Conservative Bounding Region

- First check for an intersection with a conservative bounding region
- Early reject

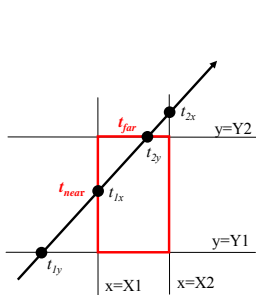


Conservative Bounding Regions

- tight → avoid false positives
- fast to intersect

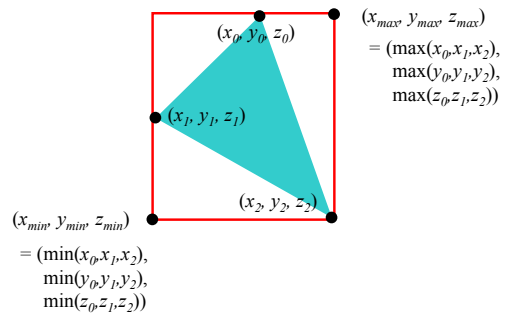


Intersection with Axis-Aligned Box

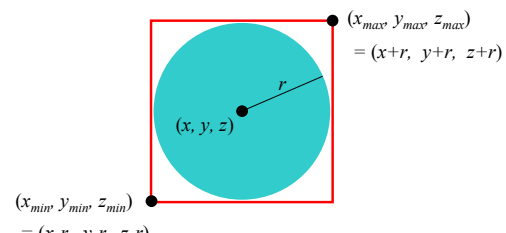


- For all 3 axes, calculate the intersection distances t_1 and t_2
- $t_{near} = \max(t_{1x}, t_{1y}, t_{1z})$
 $t_{far} = \min(t_{2x}, t_{2y}, t_{2z})$
- If $t_{near} > t_{far}$, box is missed
- If $t_{far} < t_{min}$, box is behind
- If box survived tests, report intersection at t_{near}

Bounding Box of a Triangle

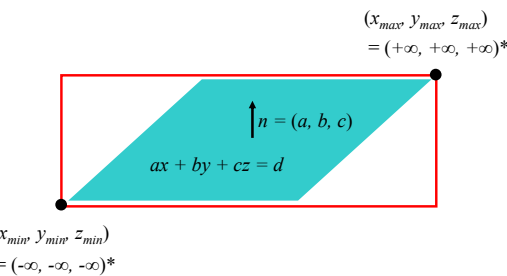


Bounding Box of a Sphere



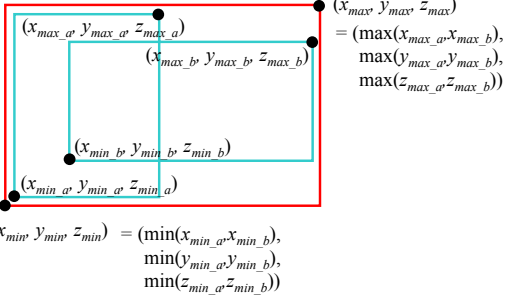
$(x_{max}, y_{max}, z_{max}) = (x+r, y+r, z+r)$
 $(x_{min}, y_{min}, z_{min}) = (x-r, y-r, z-r)$

Bounding Box of a Plane



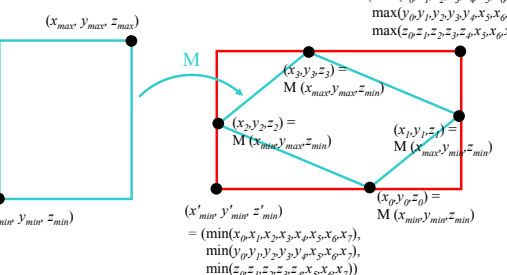
$(x_{max}, y_{max}, z_{max}) = (+\infty, +\infty, +\infty)^*$
 $(x_{min}, y_{min}, z_{min}) = (-\infty, -\infty, -\infty)^*$
** unless n is exactly perpendicular to an axis*

Bounding Box of a Group



$(x_{min}, y_{min}, z_{min}) = (\min(x_{min_a}, x_{min_b}), \min(y_{min_a}, y_{min_b}), \min(z_{min_a}, z_{min_b}))$
 $(x_{max}, y_{max}, z_{max}) = (\max(x_{max_a}, x_{max_b}), \max(y_{max_a}, y_{max_b}), \max(z_{max_a}, z_{max_b}))$

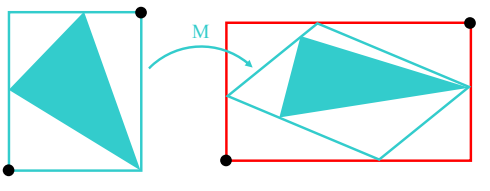
Bounding Box of a Transform



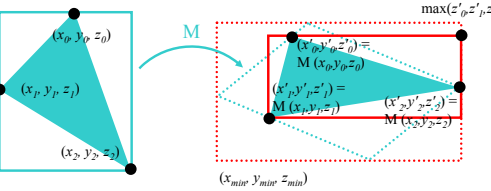
$(x'_{min}, y'_{min}, z'_{min}) = (\min(x_{p_0}, x_{p_1}, x_{p_2}, x_{p_3}, x_{p_4}, x_{p_5}, x_{p_6}, x_{p_7}), \min(y_{p_0}, y_{p_1}, y_{p_2}, y_{p_3}, y_{p_4}, y_{p_5}, y_{p_6}, y_{p_7}), \min(z_{p_0}, z_{p_1}, z_{p_2}, z_{p_3}, z_{p_4}, z_{p_5}, z_{p_6}, z_{p_7}))$
 $(x'_{max}, y'_{max}, z'_{max}) = (\max(x_{p_0}, x_{p_1}, x_{p_2}, x_{p_3}, x_{p_4}, x_{p_5}, x_{p_6}, x_{p_7}), \max(y_{p_0}, y_{p_1}, y_{p_2}, y_{p_3}, y_{p_4}, y_{p_5}, y_{p_6}, y_{p_7}), \max(z_{p_0}, z_{p_1}, z_{p_2}, z_{p_3}, z_{p_4}, z_{p_5}, z_{p_6}, z_{p_7}))$

Special Case: Transformed Triangle

Can we do better?



Special Case: Transformed Triangle

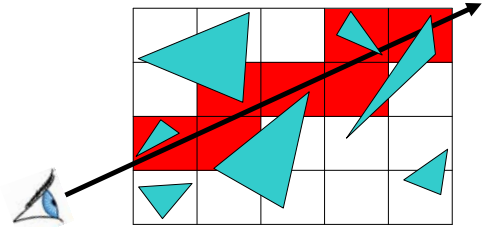


$(x'_{min}, y'_{min}, z'_{min}) = (\min(x'_{p_0}, x'_{p_1}, x'_{p_2}), \min(y'_{p_0}, y'_{p_1}, y'_{p_2}), \min(z'_{p_0}, z'_{p_1}, z'_{p_2}))$
 $(x'_{max}, y'_{max}, z'_{max}) = (\max(x'_{p_0}, x'_{p_1}, x'_{p_2}), \max(y'_{p_0}, y'_{p_1}, y'_{p_2}), \max(z'_{p_0}, z'_{p_1}, z'_{p_2}))$

Ray-tracing Acceleration

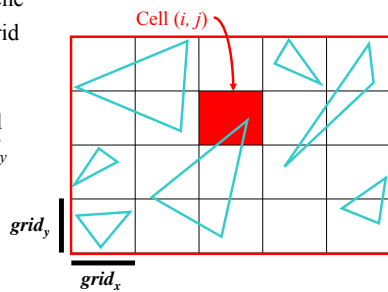
- Motivation – Distribution Ray Tracing
- Bounding Boxes
- Spatial Acceleration Data Structures
 - Regular Grid
 - Adaptive Grids
 - Hierarchical Bounding Volumes
- Flattening the Transformation Hierarchy

Regular Grid



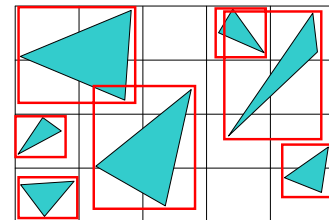
Create Grid

- Find bounding box of scene
- Choose grid resolution (n_x, n_y, n_z)
- $grid_x$ need not = $grid_y$



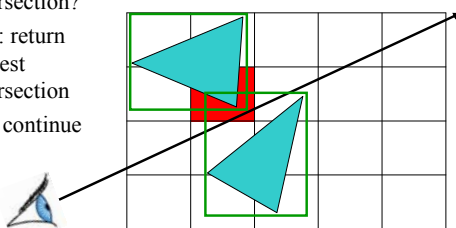
Insert Primitives into Grid

- Primitives that overlap multiple cells?
- Insert into multiple cells (use pointers)



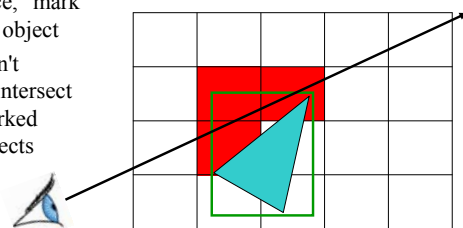
For Each Cell Along a Ray

- Does the cell contain an intersection?
- Yes: return closest intersection
- No: continue



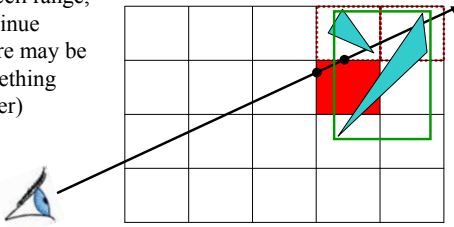
Preventing Repeated Computation

- Perform the computation once, "mark" the object
- Don't re-intersect marked objects



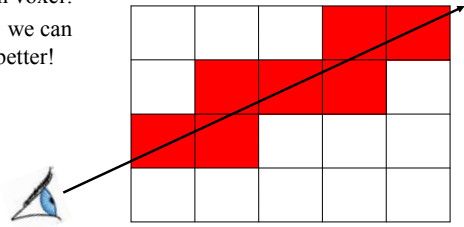
Don't Return Distant Intersections

- If intersection t is not within the cell range, continue (there may be something closer)



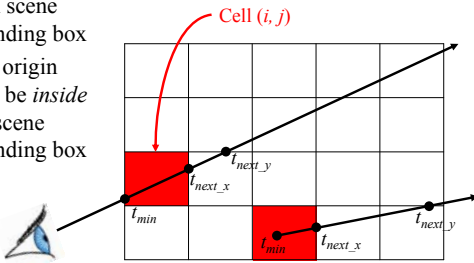
Which Cells Should We Examine?

- Should we intersect the ray with each voxel?
- No! we can do better!



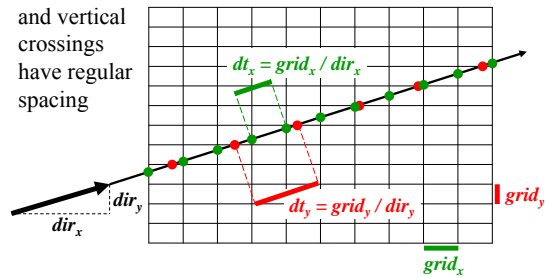
Where Do We Start?

- Intersect ray with scene bounding box
- Ray origin may be *inside* the scene bounding box



Is there a Pattern to Cell Crossings?

- Yes, the horizontal and vertical crossings have regular spacing

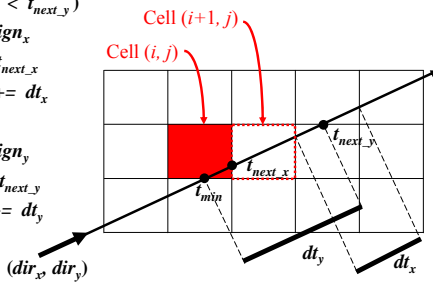


What's the Next Cell?

- ```

if (t_next_x < t_next_y)
 i += sign_x
 t_min = t_next_x
 t_next_x += dt_x
else
 j += sign_y
 t_min = t_next_y
 t_next_y += dt_y

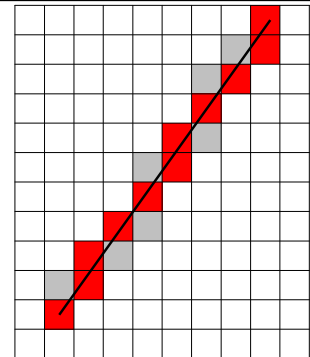
```



if ( $dir_x > 0$ )  $sign_x = 1$  else  $sign_x = -1$   
 if ( $dir_y > 0$ )  $sign_y = 1$  else  $sign_y = -1$

## What's the Next Cell?

- 3DDDA – Three Dimensional Digital Difference Analyzer
- Similar to Bresenham's Line Rasterization!



## Pseudo-Code

```

create grid
insert primitives into grid
for each ray r
 find initial cell c(i,j), tmin, tnext_x & tnext_y
 compute dtx, dty, signx and signy
 while c != NULL
 for each primitive p in c
 intersect r with p
 if intersection in range found
 return
 c = find next cell

```

## Regular Grid Discussion

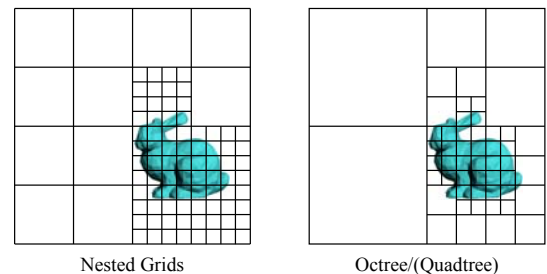
- Advantages?
  - easy to construct
  - easy to traverse
- Disadvantages?
  - may be only sparsely filled
  - geometry may still be clumped

## Ray-tracing Acceleration

- Motivation – Distribution Ray Tracing
- Bounding Boxes
- Spatial Acceleration Data Structures
  - Regular Grid
  - **Adaptive Grids**
  - Hierarchical Bounding Volumes
- Flattening the Transformation Hierarchy

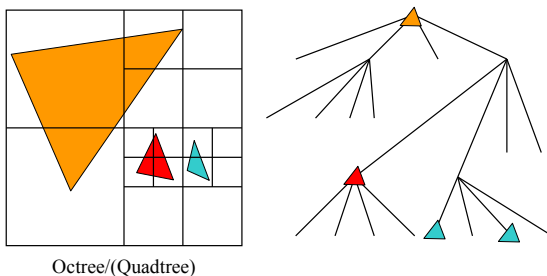
## Adaptive Grids

- Subdivide until each cell contains no more than  $n$  elements, or maximum depth  $d$  is reached



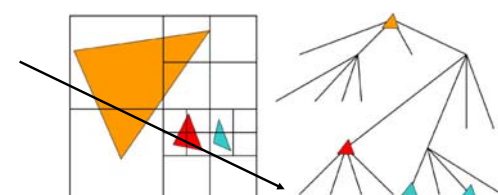
## Primitives in an Adaptive Grid

- Can live at intermediate levels, or be pushed to lowest level of grid



## Adaptive Grid Discussion

- Advantages?
  - grid complexity matches geometric density
- Disadvantages?
  - more expensive to traverse (especially octree)

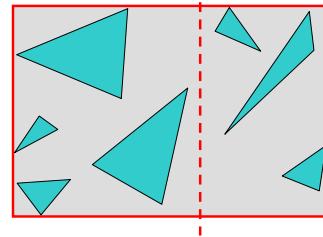


## Ray-tracing Acceleration

- Motivation – Distribution Ray Tracing
- Bounding Boxes
- Spatial Acceleration Data Structures
  - Regular Grid
  - Adaptive Grids
  - **Hierarchical Bounding Volumes**
- Flattening the Transformation Hierarchy

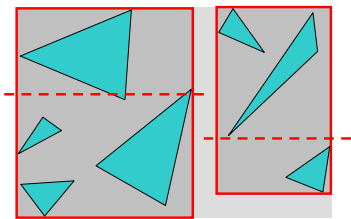
## Bounding Volume Hierarchy

- Find bounding box of objects
- Split objects into two groups
- Recurse



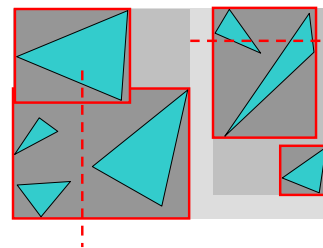
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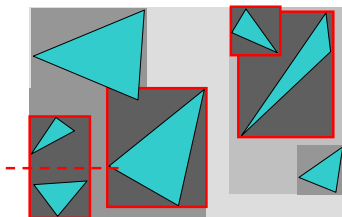
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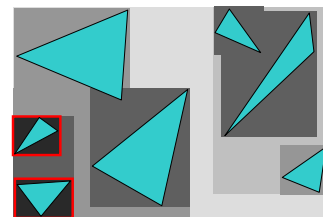
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## Bounding Volume Hierarchy

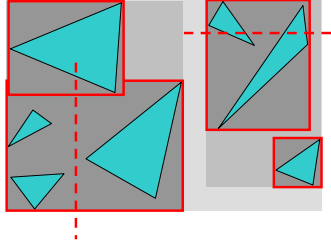
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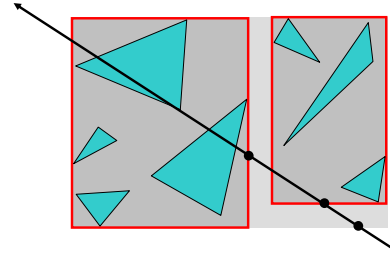
## Where to split objects?

- At midpoint *OR*
- Sort, and put half of the objects on each side *OR*
- Use modeling hierarchy



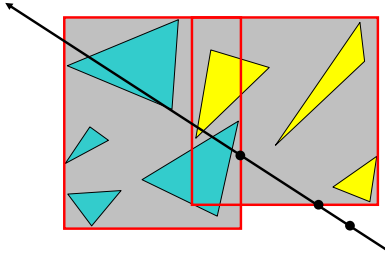
## Intersection with BVH

- Check sub-volume with closer intersection first



## Intersection with BVH

- Don't return intersection immediately if the other subvolume may have a closer intersection



## Bounding Volume Hierarchy Discussion

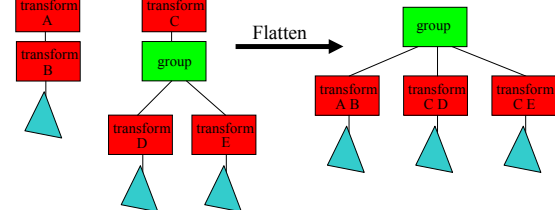
- Advantages
  - easy to construct
  - easy to traverse
  - binary
- Disadvantages
  - may be difficult to choose a good split for a node
  - poor split may result in minimal spatial pruning

## Ray-tracing Acceleration

- Motivation – Distribution Ray Tracing
- Bounding Boxes
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- **Flattening the Transformation Hierarchy**

## Transformation Hierarchy

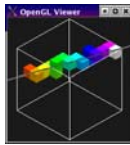
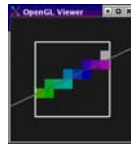
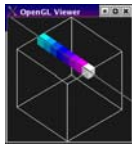
- Group & Transformation hierarchy may not be a good spatial hierarchy



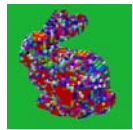
## Ray Marching Visualization



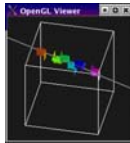
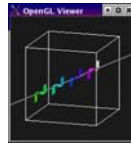
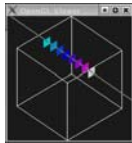
sphere voxelization



cells traversed



primitive density



entered faces

- Next time: ray-tracing at Pixar